## **End of Result Set**

Generate Collection

L7: Entry 2 of 2

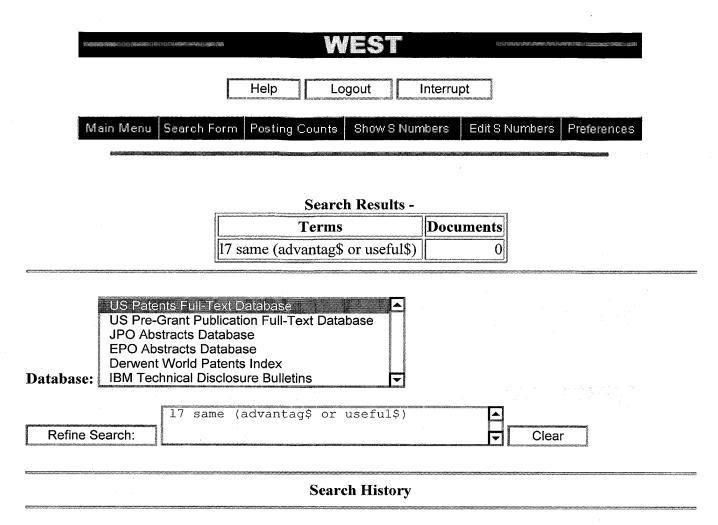
File: USPT

Sep 11, 2001

DOCUMENT-IDENTIFIER: US 6287762 B1 TITLE: Purification of a triple helix formation with an immobilized oligonucleotide

## DEPL:

The amplified DNA fragment 124 base pairs in length is separated by electrophoresis on 3% agarose gel in the presence of  $\underline{\text{SybrGreen I}}$  (Molecular Probes, Eugene, U.S.A.), and then quantified by reference to an Ultrapur genomic DNA series from E. coli strain B (Sigma, ref D4889).



Today's Date: 10/10/2001

DB Name	<u>Query</u>	Hit Count	Set Name
USPT	17 same (advantag\$ or useful\$)	0	<u>L8</u>
USPT	SybrGreen adj I	2	<u>L7</u>
USPT	SybrGreen1	0	<u>L6</u>
USPT	SybrGreenI	0	<u>L5</u>
USPT	11 same (SybrGreenI)	0	<u>L4</u>
USPT	11 same detect\$ same (SybrGreenI)	0	<u>L3</u>
USPT	11 same detect\$ same (SybrGreen near0 I)	0	<u>L2</u>
USPT	amplif\$ same (nucleic or DNA or RNA or oligo\$)	18748	<u>L1</u>

3 ANSWER 2 OF 4 CAPLUS COPYRIGHT 2001 ACS

AN 2001:32537 CAPLUS

 ${\tt TI}$  Target sequence quantification of genome DNA and mRNA with a light cycler system

AU Takahashi, Setsuko; Matsukawa, Shigeru

CS Center for Experimental Equipment, Fukui Medical University, Japan

SO Seirigaku Gijutsu Kenkyukai Hokoku (2000), 22, 59-61 CODEN: SGKHEB; ISSN: 0285-3299

PB Okazaki Kokuritsu Kyodo Kenkyu Kiko, Seirigaku Kenkyusho Gijutsuka

DT Journal

LA Japanese

AB The Light Cycler System, a fast PCR amplification and anal. system, completes 30 PCR cycles within just 30 min. It can det. the amt. of target sequences in starting materials by real time measuring of DNA-SYBRGreen I fluorescence in the PCR log-linear phase.

This report shows the optimization of exptl. procedures for

quantification

of c-myc proto-oncogene DNA present and c-myc mRNA expressing in  ${\rm HL}60$  and  ${\rm K5}62~{\rm h}$